

**NOTICE TO COMPLY WITH REQUIREMENTS FOR PATENT APPLICATIONS CONTAINING NUCLEOTIDE SEQUENCE AND/OR AMINO ACID SEQUENCE DISCLOSURES**

The nucleotide and/or amino acid sequence disclosure contained in this application does not comply with the requirements for such a disclosure as set forth in 37 CFR 1.821 - 1.825 for the following reason(s):

- ☐ 1. This application clearly fails to comply with the requirements of 37 CFR 1.821 - 1.825. Applicant's attention is directed to these regulations, published at 1114 OG 29, May 15, 1990 and at 55 FR 18230, May 1, 1990.
- ☐ 2. This application does not contain, as a separate part of the disclosure on paper copy, a "Sequence Listing" as required by 37 CFR 1.821(c).
- ☒ 3. A copy of the "Sequence Listing" in computer readable form has not been submitted as required by 37 CFR 1.821(e).
- ☐ 4. A copy of the "Sequence Listing" in computer readable form has been submitted. However, the content of the computer readable form does not comply with the requirements of 37 CFR 1.822 and/or 1.823, as indicated on the attached marked-up copy of the "Raw Sequence Listing."
- ☐ 5. The computer readable form that has been filed with this application has been found to be damaged and/or unreadable as indicated on the attached CRF Diskette Problem Report. A substitute computer readable form must be submitted as required by 37 CFR 1.825(d).
- ☐ 6. The paper copy of the "Sequence Listing" is not the same as the computer readable form of the "Sequence Listing" as required by 37 CFR 1.821(e).
- ☐ 7. Other: \_\_\_\_\_

**Applicant must provide:**

- ☐ An initial or substitute computer readable form (CRF) copy of the "Sequence Listing"
- ☐ An initial or substitute paper copy of the "Sequence Listing", as well as an amendment directing its entry into the specification
- ☒ A statement that the content of the paper and computer readable copies are the same and, where applicable, include no new matter, as required by 37 CFR 1.821(e) or 1.821(f) or 1.821(g) or 1.825(b) or 1.825(d)

For questions regarding compliance with these requirements, please contact:

For Rules Interpretation, call (703) 308-1123  
For CRF submission help, call (703) 308-4212  
For PatentIn software help, call (703) 308-6856

Please return a copy of this notice with your response.

RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/487,461DATE: 09/26/95  
TIME: 15:55:05

INPUT SET: S6366.raw

This Raw Listing contains the General  
Information Section and up to the first 5 pages.

## SEQUENCE LISTING

## (1) General Information:

- (i) APPLICANT: Parker, Thomas S.; Levine, Daniel M.;  
Rubin, Albert L; Gordon, Bruce R; Saal, Stuart D.
- (ii) TITLE OF INVENTION: Methods Useful in Endotoxin Based  
Prophylaxis and Therapy
- (iii) NUMBER OF SEQUENCES: 9
- (iv) CORRESPONDENCE ADDRESS:  
(A) ADDRESSEE: Felfe & Lynch  
(B) STREET: 805 Third Avenue  
(C) CITY: New York City  
(D) STATE: New York  
(E) COUNTRY: USA  
(F) ZIP: 10022
- (v) COMPUTER READABLE FORM:  
(A) MEDIUM TYPE: Diskette, 5.25 inch, 360 kb  
storage  
(B) COMPUTER: IBM PS/2  
(C) OPERATING SYSTEM: PC-DOS  
(D) SOFTWARE: Wordperfect
- (vi) CURRENT APPLICATION DATA:  
(A) APPLICATION NUMBER: 08/487,461  
(B) FILING DATE: June 7, 1995  
(C) CLASSIFICATION: 435
- (vii) PRIOR APPLICATION DATA:  
(A) APPLICATION NUMBER: PCT/US93/07453  
(B) FILING DATE: 9-AUGUST-1993
- (vii) PRIOR APPLICATION DATA:  
(A) APPLICATION NUMBER: 07/928,930  
(B) FILING DATE: 12-AUGUST-1992
- (viii) ATTORNEY/AGENT INFORMATION:  
(A) NAME: Hanson, Norman D.  
(B) REGISTRATION NUMBER: 30,946  
(C) REFERENCE/DOCKET NUMBER: ROGO 211.2
- (ix) TELECOMMUNICATION INFORMATION:

ENTERED

--&gt;

RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/487,461DATE: 09/26/95  
TIME: 15:55:08

INPUT SET: S6366.raw

47 (A) TELEPHONE: (212) 688-9200  
48 (B) TELEFAX: (212) 838-3884  
49  
50 (2) INFORMATION FOR SEQ ID NO: 1:  
51 (i) SEQUENCE CHARACTERISTICS:  
52 (A) LENGTH: 18 amino acids  
53 (B) TYPE: amino acid  
54 (D) TOPOLOGY: linear  
55 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:  
56  
57 Asp Trp Leu Lys Ala Phe Tyr Asp Lys Val Ala Glu Lys Leu Lys  
58 5 10 15  
59 Glu Ala Phe  
60  
61  
62  
63  
64 (2) INFORMATION FOR SEQ ID NO: 2:  
65 (i) SEQUENCE CHARACTERISTICS:  
66 (A) LENGTH: 18 amino acids  
67 (B) TYPE: amino acid  
68 (D) TOPOLOGY: linear  
69 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:  
70  
71 Lys Trp Leu Asp Ala Phe Tyr Lys Asp Val Ala Lys Glu Leu Glu  
72 5 10 15  
73 Lys Ala Phe  
74  
75  
76  
77  
78 (2) INFORMATION FOR SEQ ID NO: 3:  
79 (i) SEQUENCE CHARACTERISTICS:  
80 (A) LENGTH: 17 amino acids  
81 (B) TYPE: amino acid  
82 (D) TOPOLOGY: linear  
83 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:  
84  
85  
86 Asp Trp Leu Lys Ala Phe Tyr Asp Lys Ala Glu Lys Leu Lys Glu  
87 5 10 15  
88 Ala Phe  
89  
90  
91  
92  
93 (2) INFORMATION FOR SEQ ID NO: 4:  
94 (i) SEQUENCE CHARACTERISTICS:  
95 (A) LENGTH: 22 amino acids  
96 (B) TYPE: amino acid  
97 (D) TOPOLOGY: linear  
98 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:  
99

RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/487,461DATE: 09/26/95  
TIME: 15:55:11

INPUT SET: S6366.raw

100 Pro Lys Leu Glu Glu Leu Lys Glu Lys Leu Lys Glu Leu Leu Glu  
101                   5                   10                   15

102 Lys Leu Lys Glu Lys Leu Ala  
103                   20

104

105

106

107 (2) INFORMATION FOR SEQ ID NO: 5:

108 (i) SEQUENCE CHARACTERISTICS:

109 (A) LENGTH: 16 amino acids

110 (B) TYPE: amino acid

111 (D) TOPOLOGY: linear

112 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:

113

114 Val Ser Ser Leu Lys Glu Tyr Trp Ser Ser Leu Lys Glu Ser Phe  
115                   5                   10                   15

116 Ser

117

118

119

120 (2) INFORMATION FOR SEQ ID NO: 6:

121 (i) SEQUENCE CHARACTERISTICS:

122 (A) LENGTH: 20 amino acids

123 (B) TYPE: amino acid

124 (D) TOPOLOGY: linear

125 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:

126

127 Val Ser Ser Leu Leu Ser Ser Leu Lys Glu Tyr Trp Ser Ser Leu  
128                   5                   10                   15

129 Lys Glu Ser Leu Ser

130                   20

131

132

133 (2) INFORMATION FOR SEQ ID NO: 7:

134 (i) SEQUENCE CHARACTERISTICS:

135 (A) LENGTH: 24 amino acids

136 (B) TYPE: amino acid

137 (D) TOPOLOGY: linear

138 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7:

139

140 Val Ser Ser Leu Leu Ser Ser Leu Leu Ser Ser Leu Lys Glu Tyr  
141                   5                   10                   15

142 Trp Ser Ser Leu Lys Glu Ser Glu Ser

143                   20

144

145

146

147 (2) INFORMATION FOR SEQ ID NO: 8:

148 (i) SEQUENCE CHARACTERISTICS:

149 (A) LENGTH: 22 amino acids

150 (B) TYPE: amino acid

151 (D) TOPOLOGY: linear

152 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 8:

RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/487,461DATE: 09/26/95  
TIME: 15:55:13

INPUT SET: S6366.raw

153  
154 Pro Val Leu Asp Glu Phe Arg Glu Lys Leu Asn Glu Glu Leu Glu  
155 5 10 15  
156 Ala Leu Lys Gln Lys Met Lys  
157 20  
158  
159  
160  
161 (2) INFORMATION FOR SEQ ID NO: 9:  
162 (i) SEQUENCE CHARACTERISTICS:  
163 (A) LENGTH: 22 amino acids  
164 (B) TYPE: amino acid  
165 (D) TOPOLOGY: linear  
166 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 9:  
167  
168 Pro Leu Ala Glu Asp Leu Gln Thr Lys Leu Asn Glu Asn Val Glu  
169 5 10 15  
170 Asp Leu Arg Lys Gln Leu Val  
171 20  
172